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Figure 1

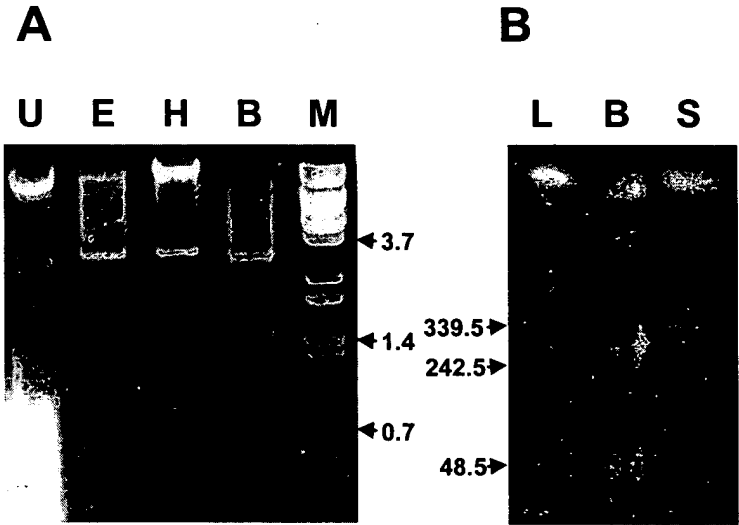


Figure 2

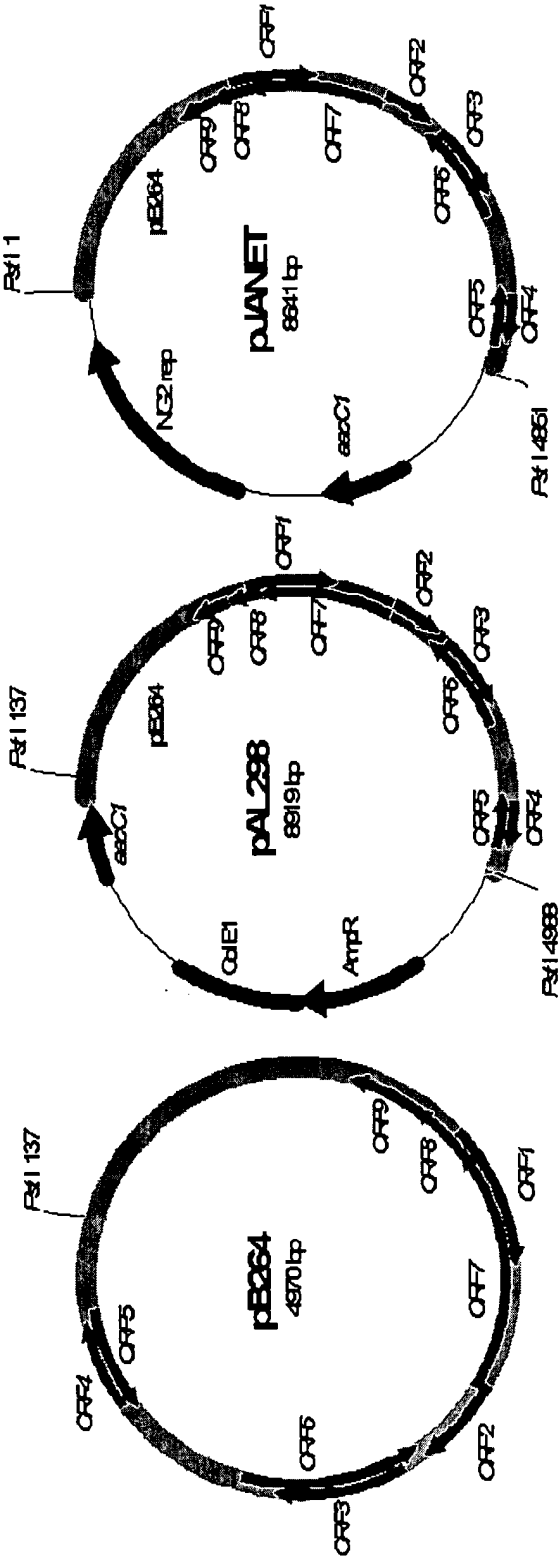


Figure 3

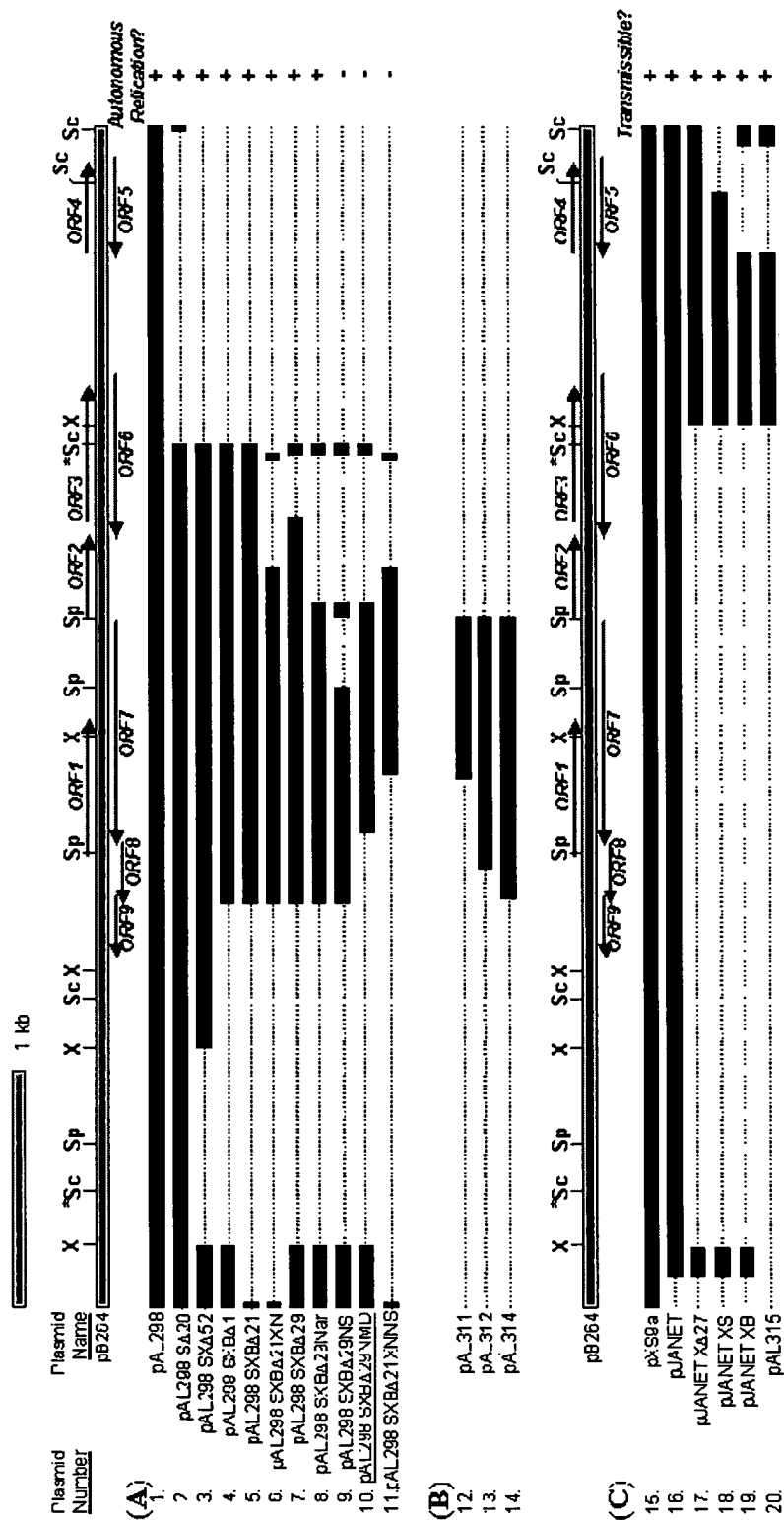


Figure 4

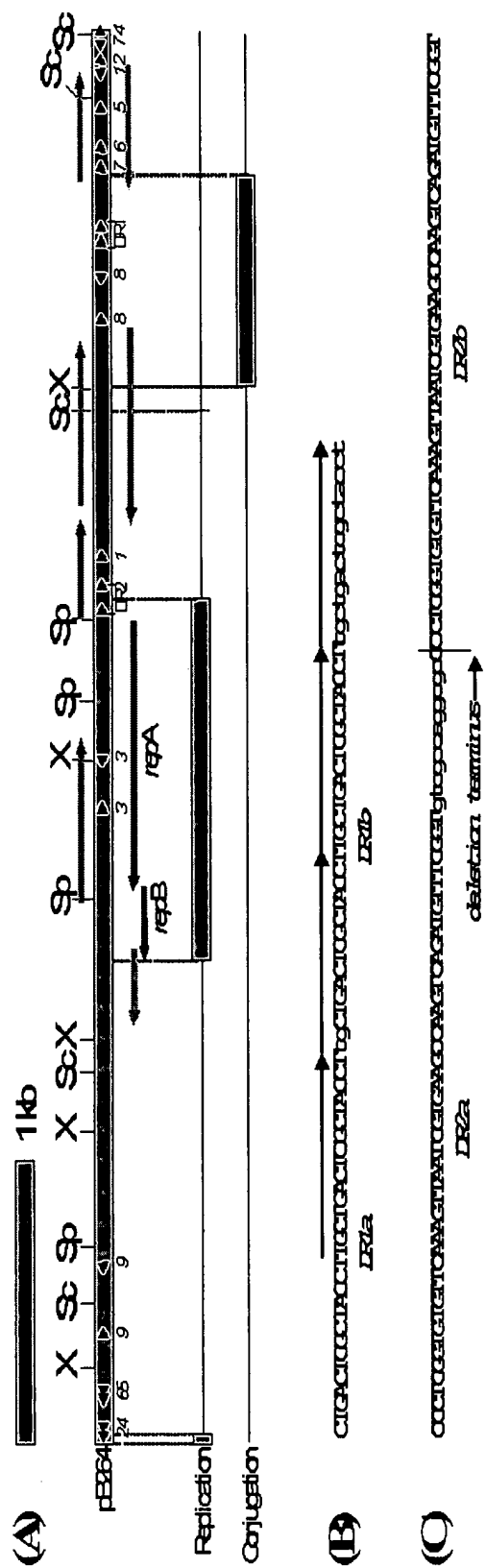


Figure 5

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GGGTGCTCCGGAGGCTCACGGCCCCGGTGGCCGATCTGCTCGCCCAGATCCACT
CGAGTGCAGGCACCAACGCGTACATCGACCCGGCGCAGGCCGTGCAGGCAGTGC
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GCGTGACGTTTACCACGGTGCGGGCGAGATCAGCGATCTTGACGTTTCATGTTCT
GTGAGGTTTCGCCTGAGAATGTCGAAGGCGTCCTCAGACGACACGTCGTGCAGTG
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TCACTGAAGAGATTCAGTGCTCCAAC TTTGTGGCCGTCGACGAGCAGACGAAAT
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GCGGCCGGCCGCGGAAGTGGTGCTCGGACGAGTG (SEQ ID NO: 1)

Figure 6

CGTCATCGCAGCTGATACTCATCGCTCGATTCCCTGCCAGGGCGTGATGGACAGTG
CCGACCGAGCATCCCAGCTCATTGGCGATAGCTCTCATGGACATGCCAGTACCG
CGTAGCTCCCGTATCTTCTCGCGCCGTTGCTGTGCCCGGGACAGGTAGGCCTCTC
GCGGCTCTGATGTCCACCGGATGATGCTGGCCGTCGATACGCCTGTGCGTTTCGGC
GAGCTCCCGCACTGTTATCCCGTTGCGAGGCAGACGCTCAGTCATGTGTGGCCTC
CTCAATTGCCAGTCGCACCTGATCCAGTTTGGTGCTGCGTGACGCCGGGCCTGG
CCGCTGGCGACGCCGCCTTTGCGGCCACGGGCGGATTGGATGGTCGAGAGTGTG
GCTTCGTAGACGGCGATAACCGTCTTTCCAGATGCGGGATCGCGTGGTGATCCATC
GGTGAATGCTGGCGGCGATGGCGCGGGCTTCAGAGGGTGGGAGTTCCTCGGTGT
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Figure 7

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TAAATCCCGGGCCACGCAACATCAGGCTTGCGTGGTACAGCCGGGAACCAGCAG
GGTGGCGGCGACACAGACCGCCGCGTAAACCTCGGACGGCTCGGAACCGTTGTA
GACAATGTCGGCGAGCGCGGCGAATACGGTCCCCGGATCCGACTTGCTGGGGGT
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CGTCAAACGCTACGCGGTGGAACCTGCTCGGGAGTCATCCAGATGGTGGAGTGA
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GGACAGCACGCGGCGCAGCCTGACTCGCTACCTTGCTGACTCGCTACCTTGCTGA
CTCGCTACCTTGCTGACTCGCTACCTTGCTGACTCGCTACCTACGCGGCGCAGCA
AGATGGAGACTTAGGTGCCCTAAGTCGCAGCCCGCATTGCGGGGATCCCTGTCG
GGGCGCTGCGCTGGGCGGTGAAATACCAACCTTGCGGTTCCCTGGGAACAGCTAA
ACCGAACGGTGTGCTTCGGCAGTCGGGGATGACGACGCC (SEQ ID NO: 3)

Figure 8

GGCGCCTGGCGACACCGAAACATCTGACTTGGCTTCACGATTAAC TTTGAACAC
ACCgAGGgAAATGCCCAGGTTTCGTTCGAGAGCATGC (SEQ ID NO:4)

Figure 9

ATGGACACACACGGGGTGCGGGAAGGTGGGGACTGGGAGCAAATGTGGCTACC
GTTcTGGCCTTTGGCAACAAACGACTTCCTTGAGGGGGTcTATCGAATGCGCCGC
CCTGCCGCACTTGAGCGGCGCTACATCGAGGCCAACCCGCAGACACTGAGCAAT
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CAGGCAATCACCTATGCCGAACGCCGTGATCGAAAACAGCAGCAACGGGCATG
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CACCACGCGATCCCGCATCTGGAAAGACGGTATCGCCGTCTACGAAGCCACACT
CTCGACCATCCAATCCGCCCCGTGGCCGCAAAGGCGGCGTCGCCAGCGGCCAGGC
CCGGCGTGACGCAGCACCAAACCTGGATCAGGTGCGACTGGCAATTGAGGAGGC
CACACATGACTGA (SEQ ID NO:5)

Figure 10

ATGACTGAGCGTCTGCCTCGCAACGGGATAACAGTGCGGGAGCTCGCCGAACGC
ACAGGCGTATCGACGGCCAGCATCATCCGGTGGACATCAGAGCCGCGAGAGGCC
TACCTGTCCCGGGCACAGCAACGGCGCGAGAAGATACGGGAGCTACGCGGTACT
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CATCACGCCCTGGCAGGAATCGAGCGATGA (SEQ ID NO:6)

Figure 11

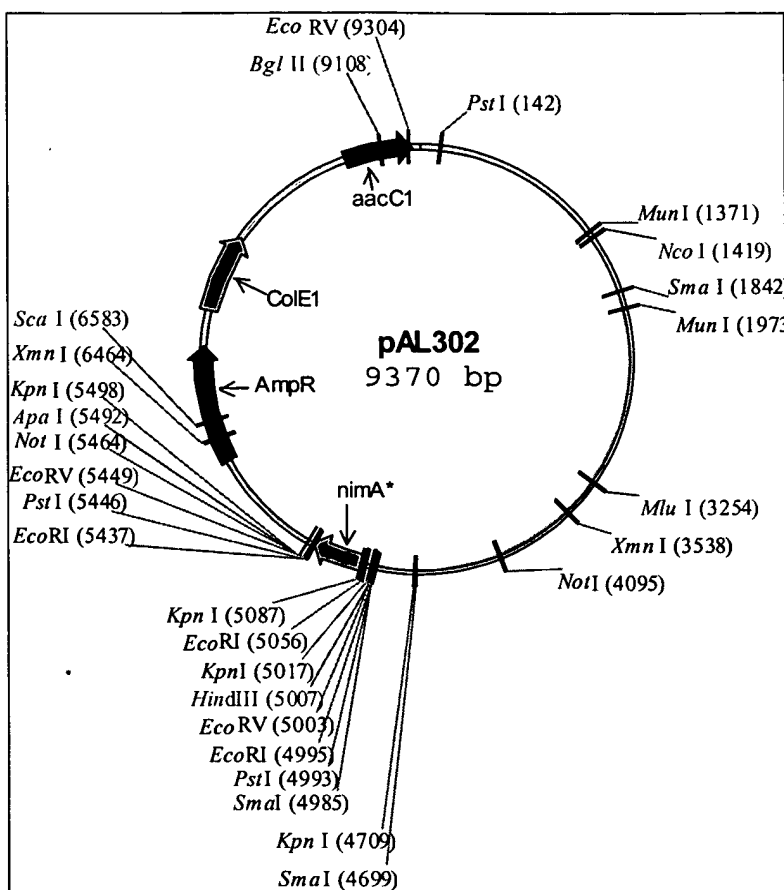


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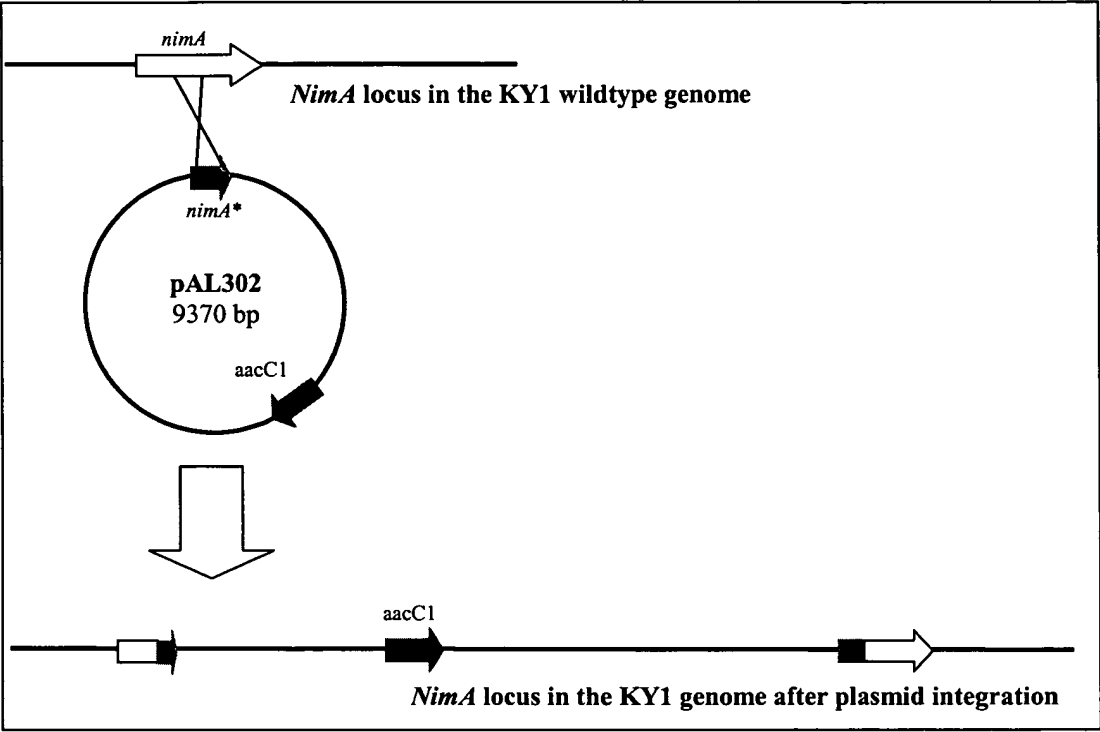


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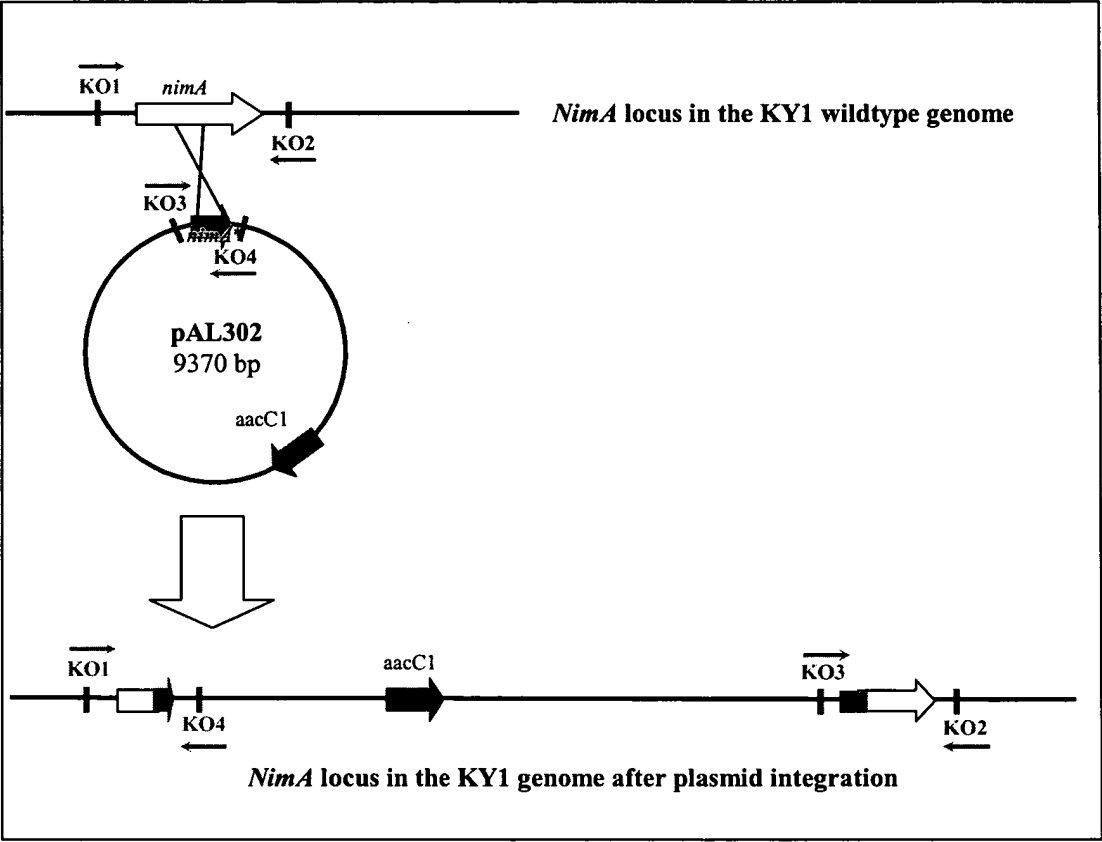


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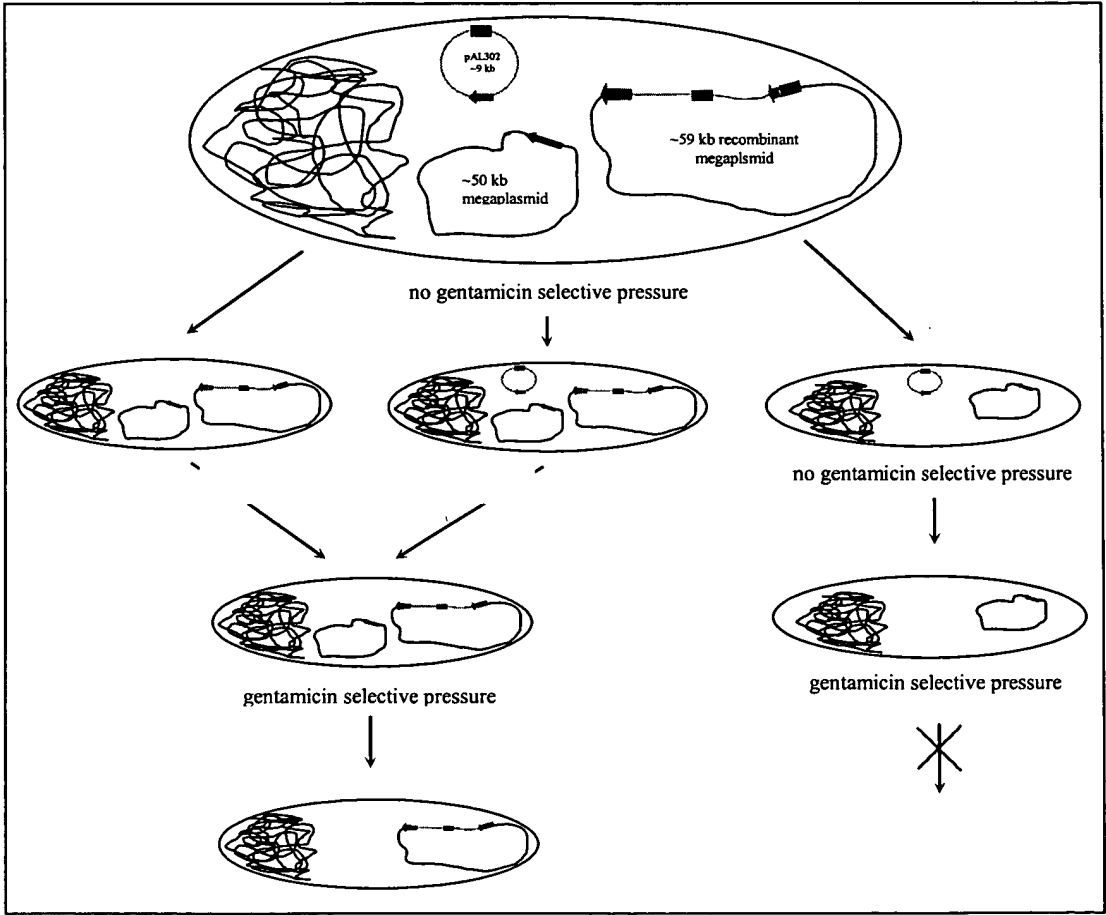


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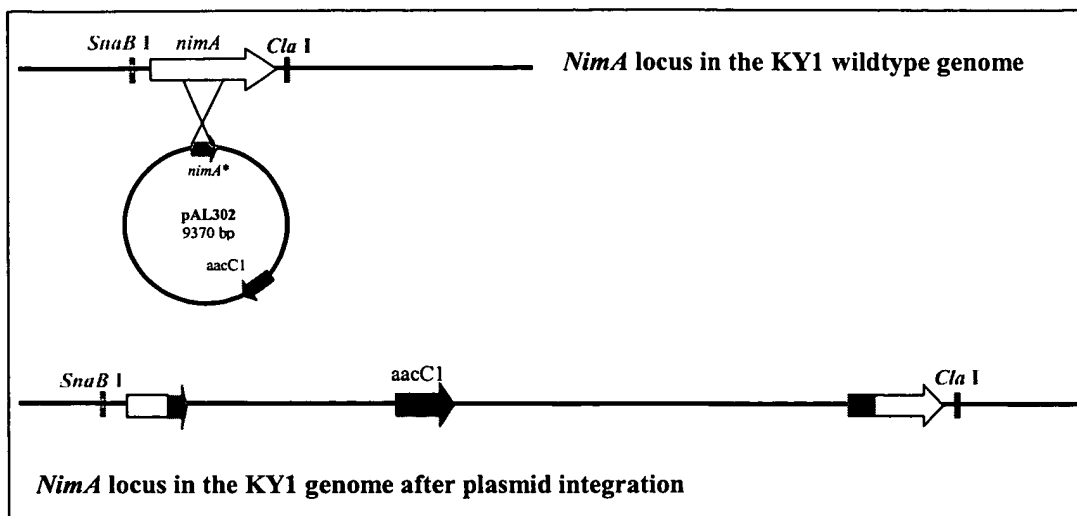


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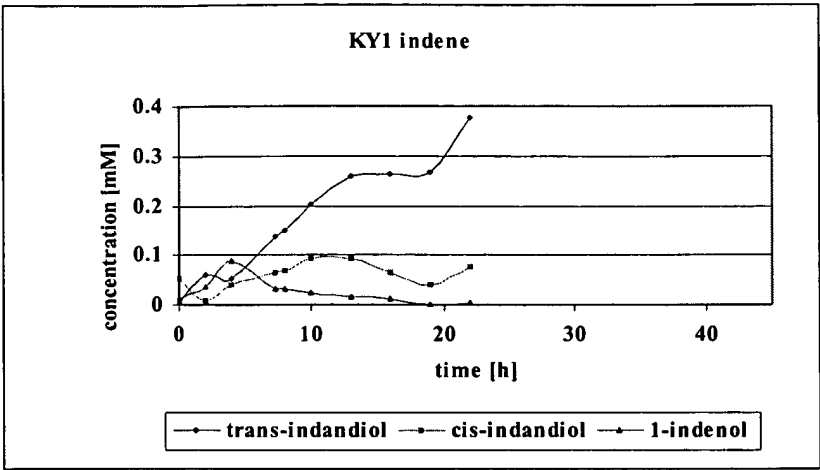


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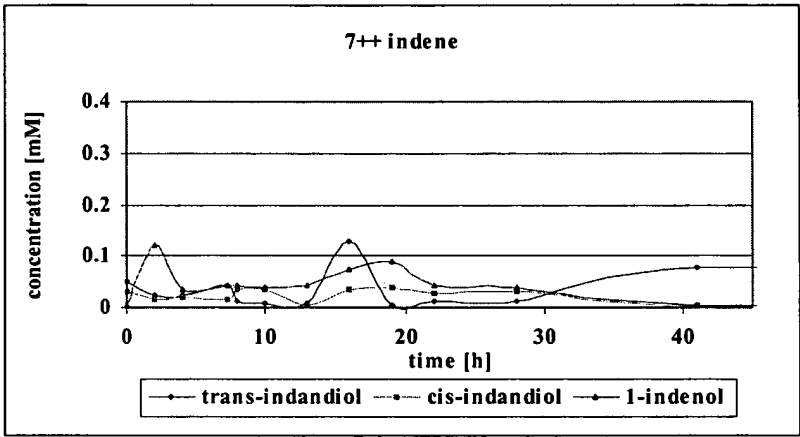


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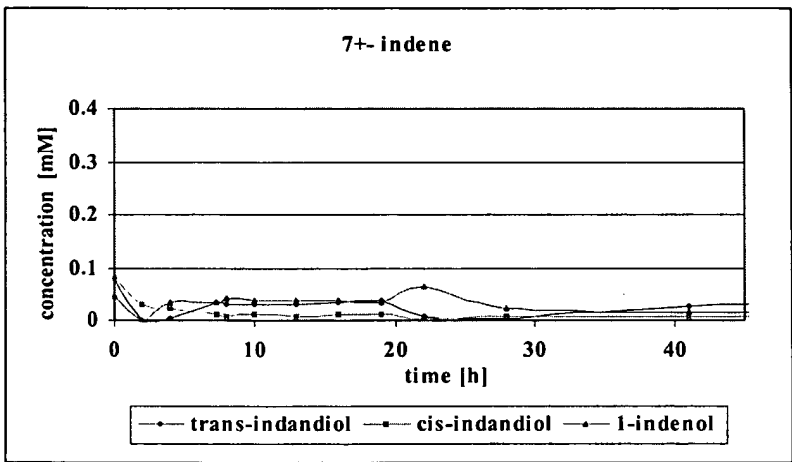


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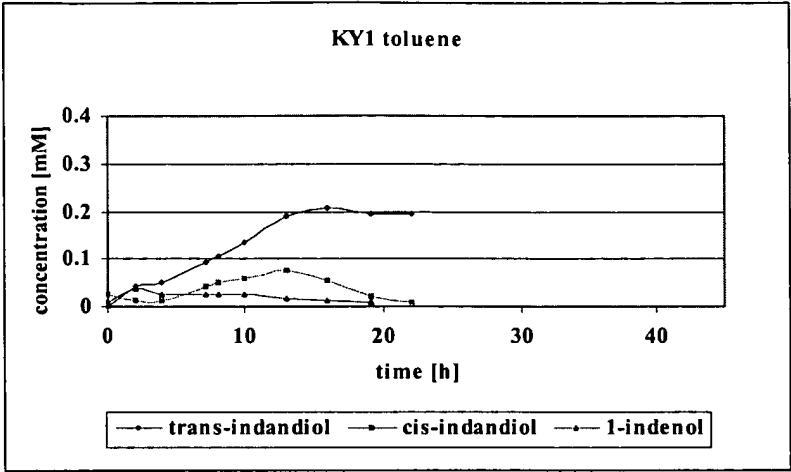


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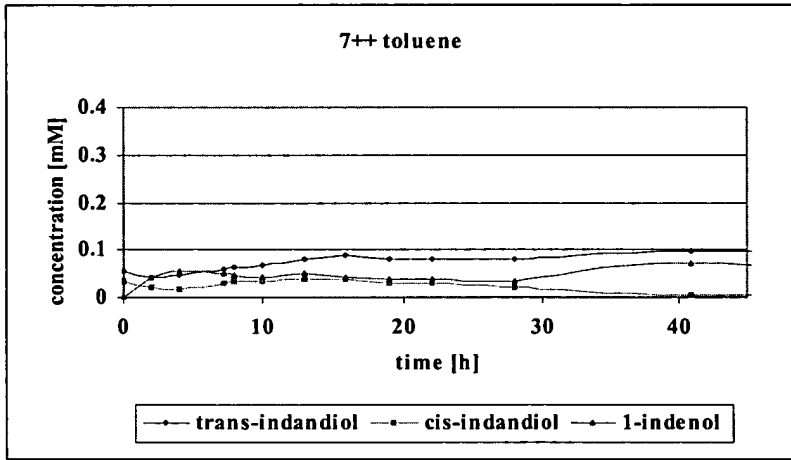


Figure 21

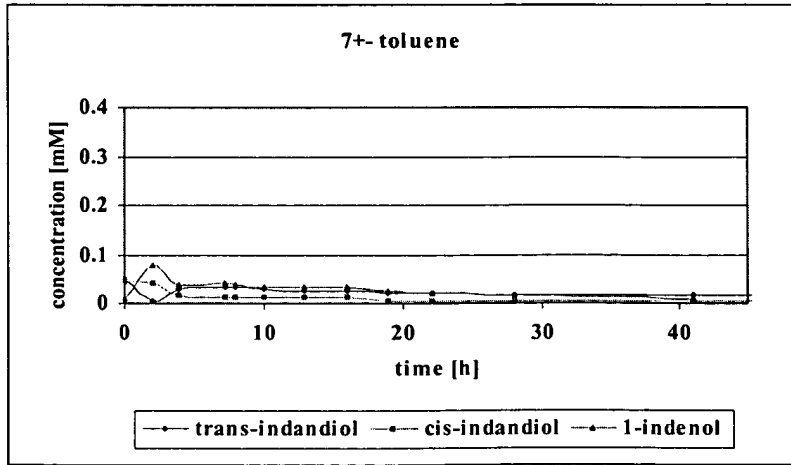


Figure 22

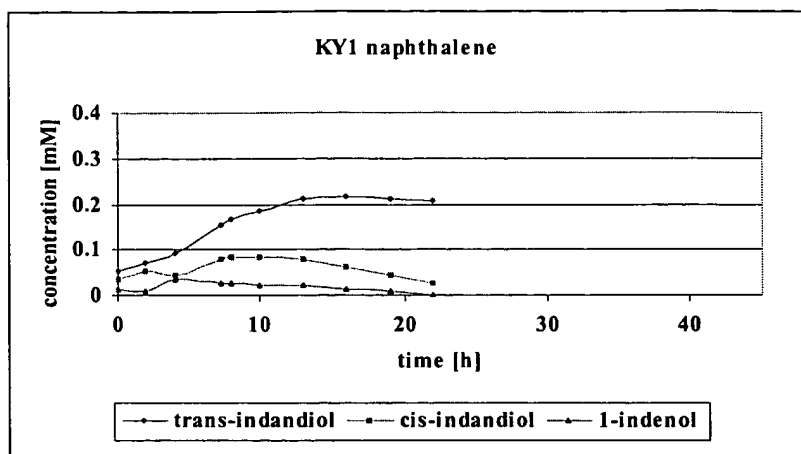


Figure 23

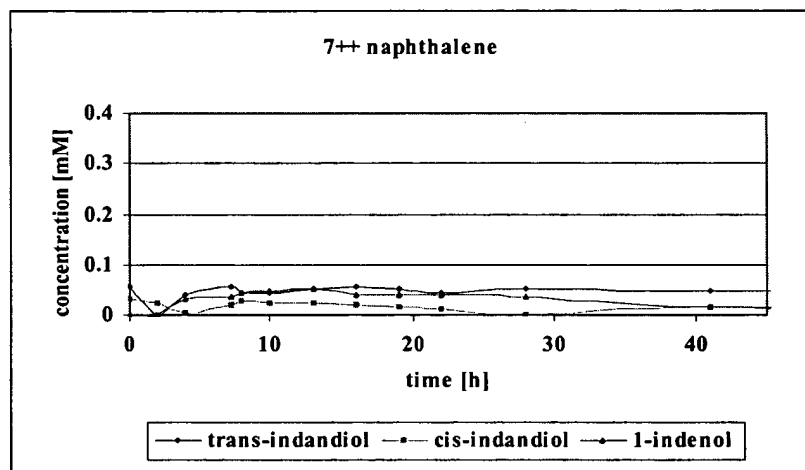


Figure 24

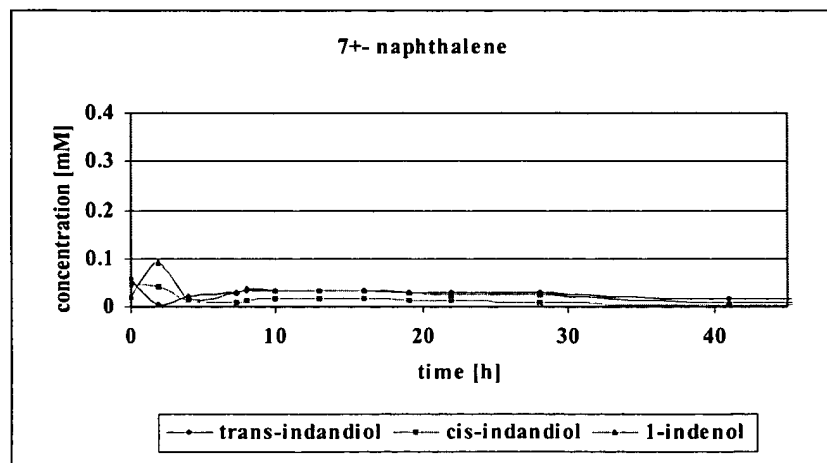


Figure 25

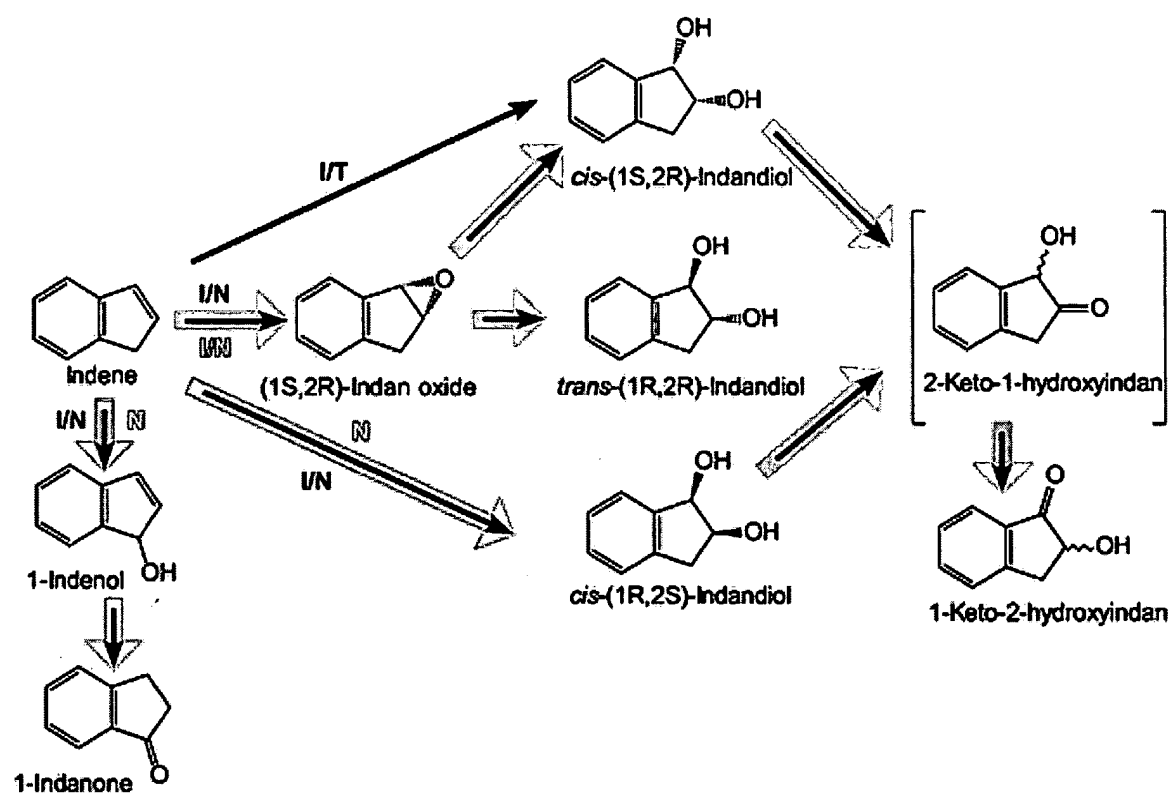


Figure 26

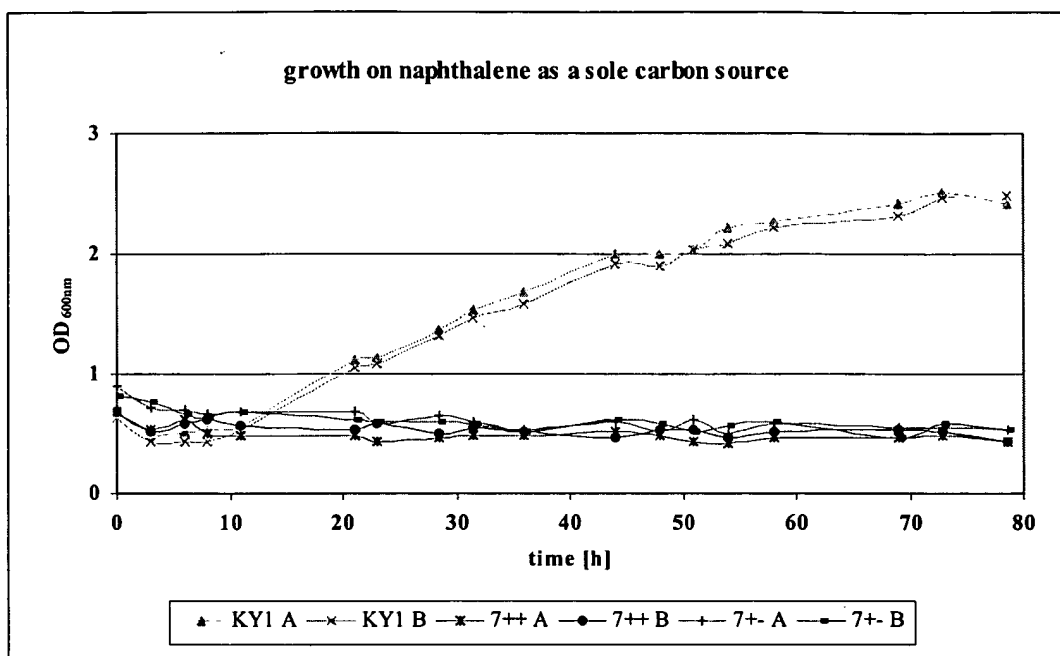
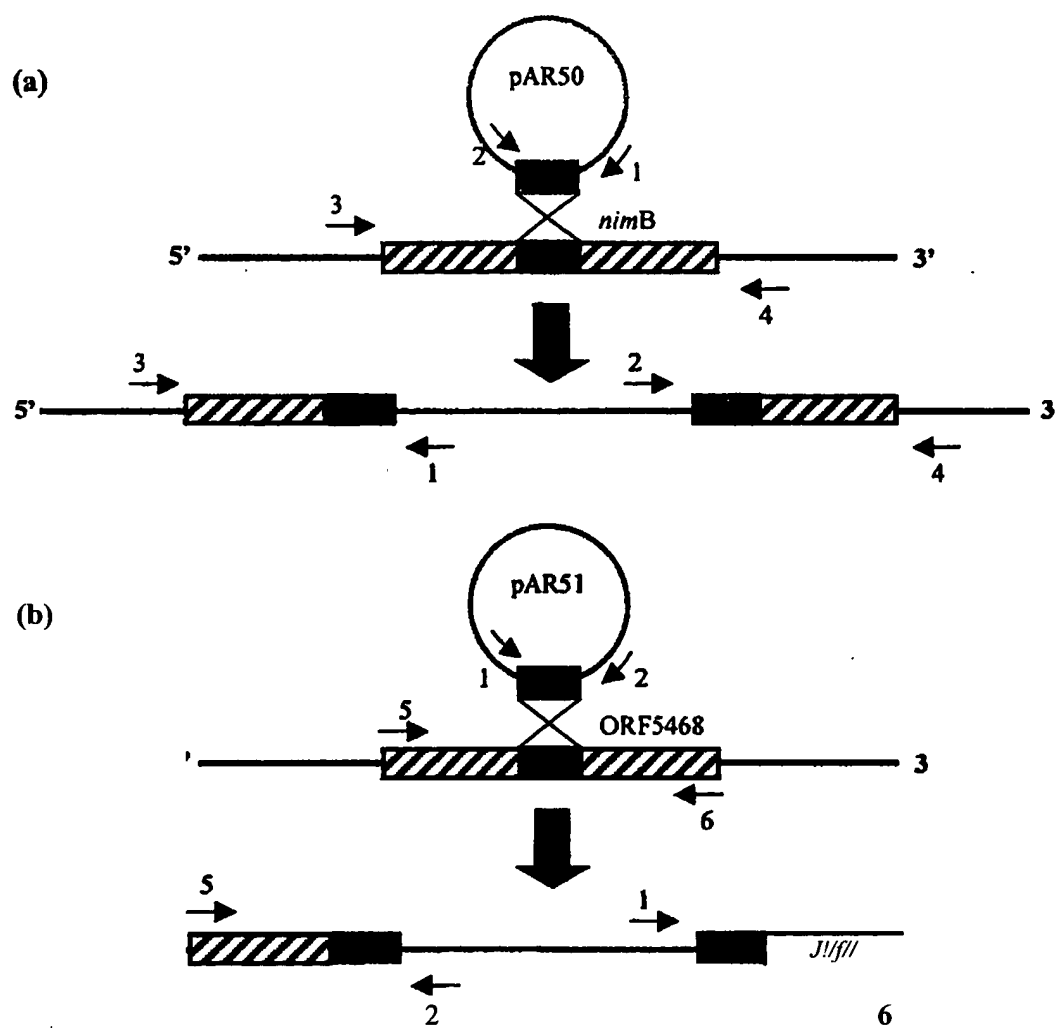


Figure 27



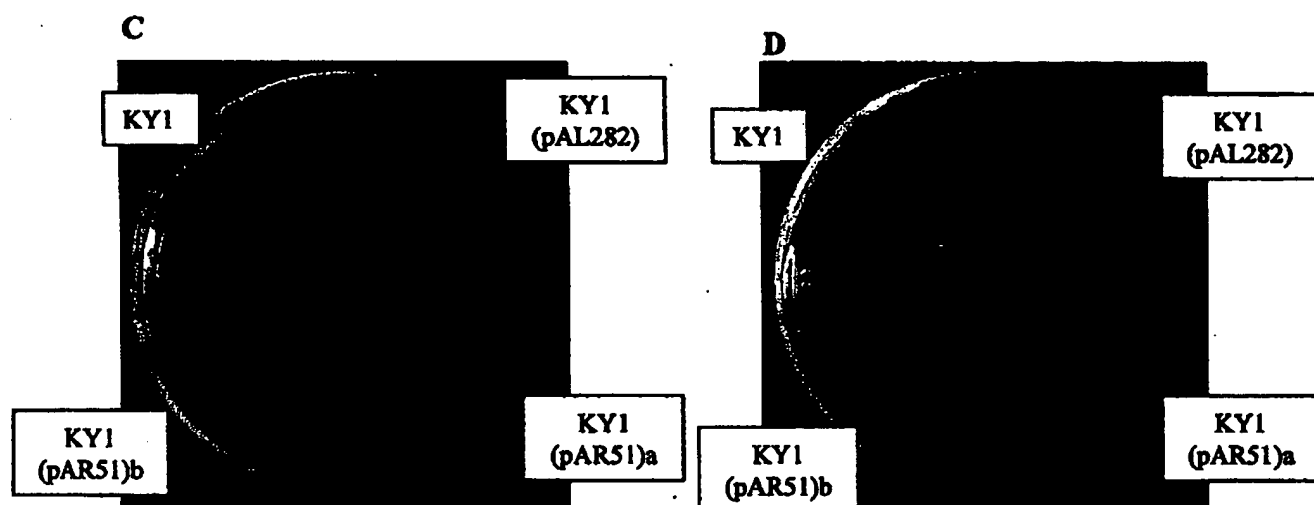
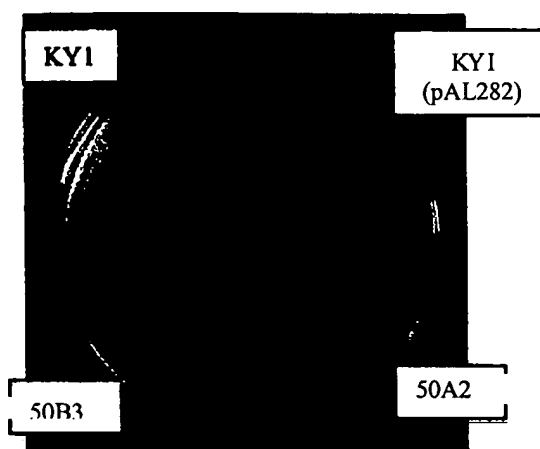
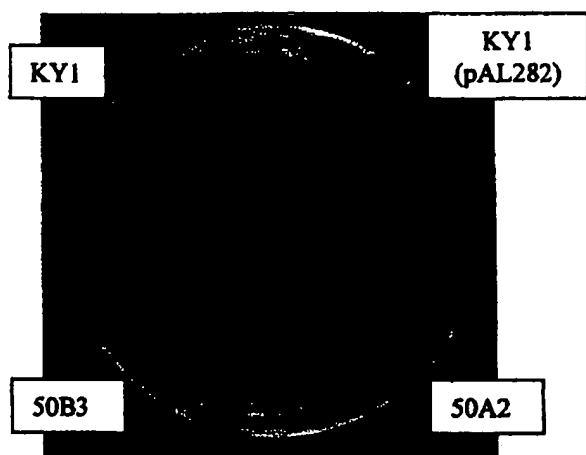


Figure 28